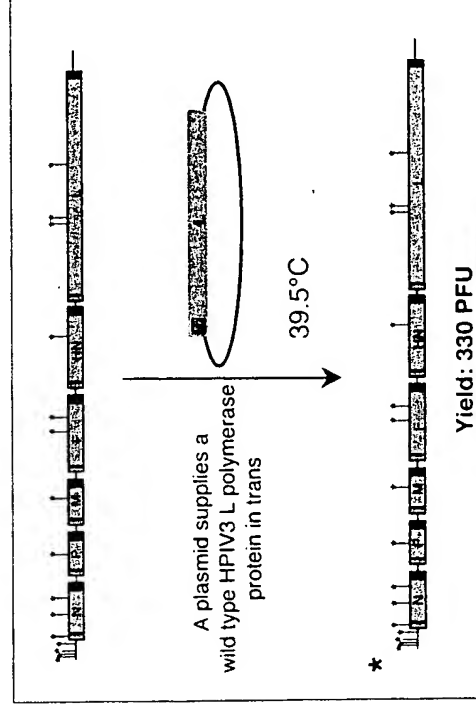
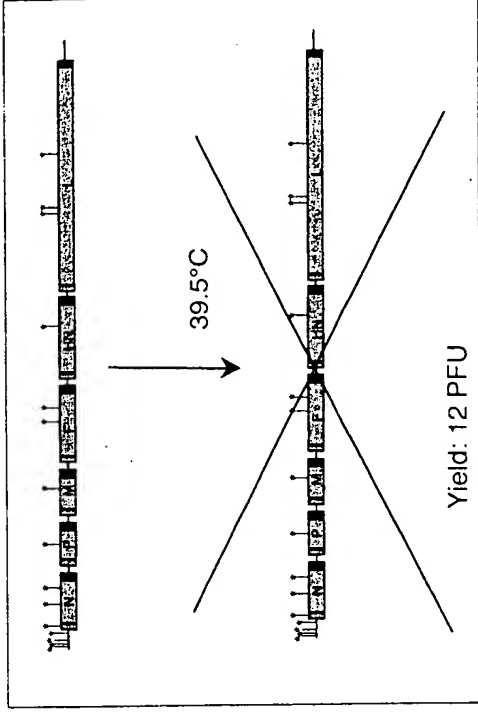
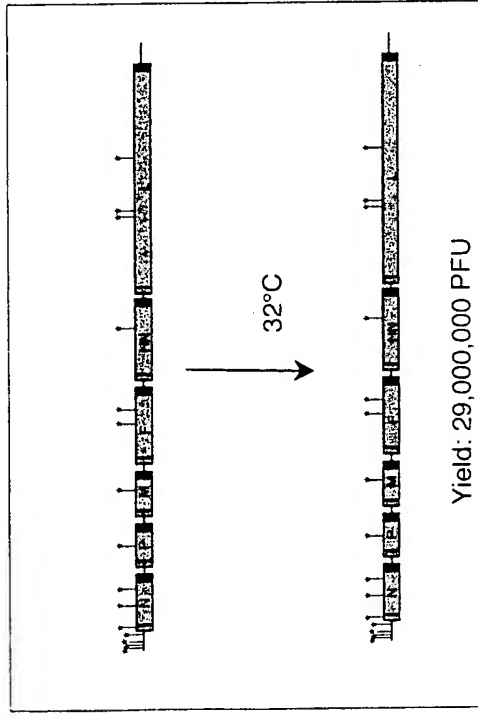


Complementation assay described in Belshe patent (No. 5,869,036) used to demonstrate that ts phenotype is due to mutations in the L polymerase



Note: wild type HPIV3 yield at 39.5 °C:  
8,000,000 PFU

Virus is unchanged\*

\* No genetic manipulation has been performed

